

Effector: Target Ratio

Fig. 1

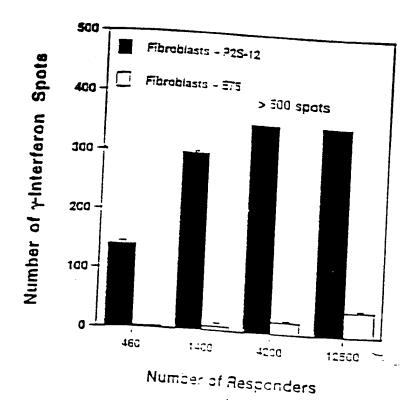


Fig. 2A

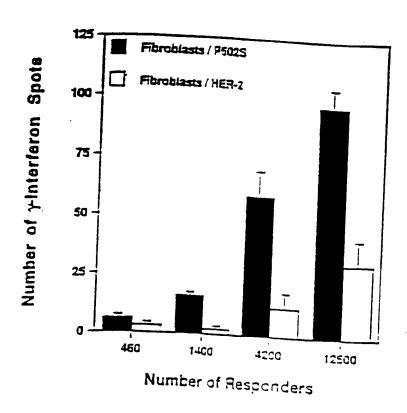


Fig. 2B

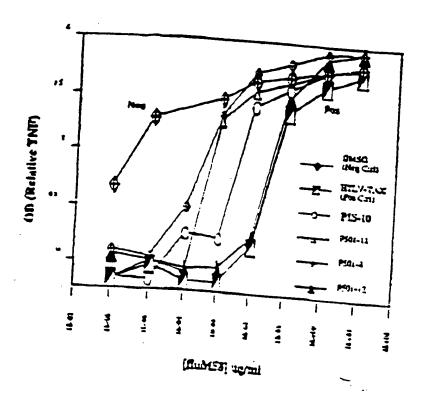


Fig. 3

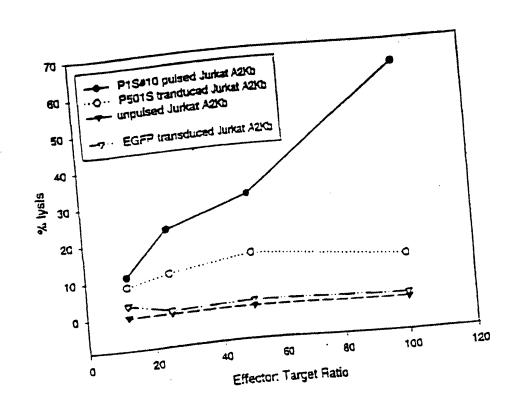


Fig. 4

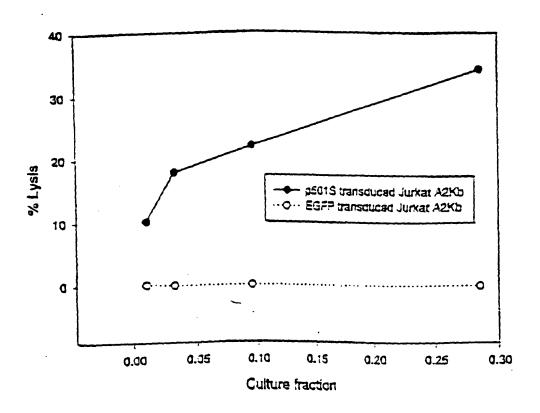
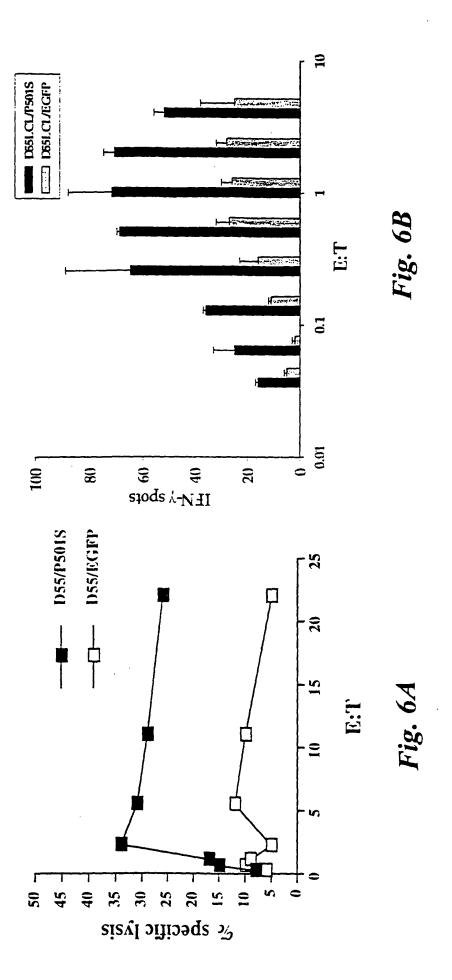
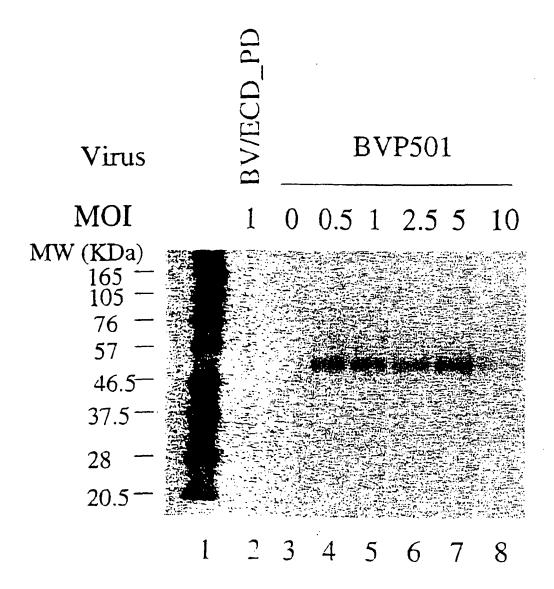


Fig. 5

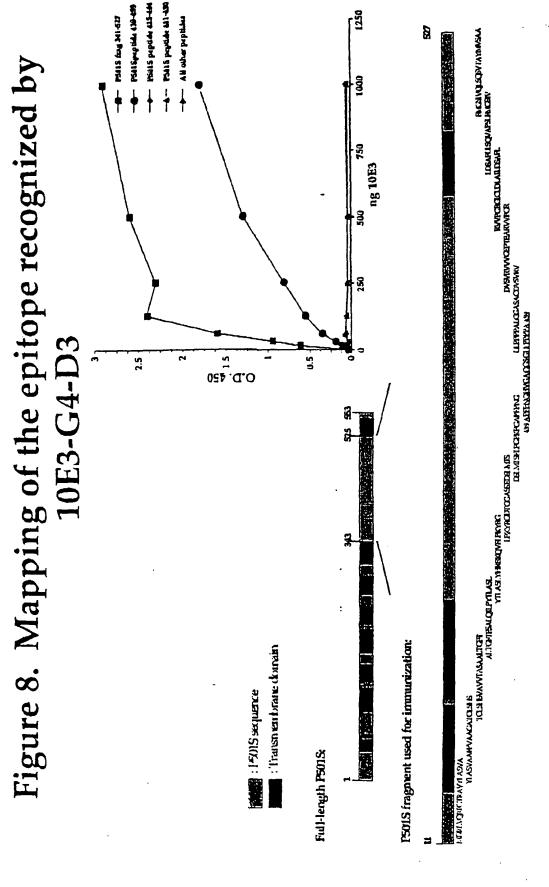


10010940.120501

Expression of P501S by the Baculovirus Expression System



0.6 million high 5 cells in 6-well plate were infected with an unrelated control virus BV/ECD_PD (lane 2), without virus (lane 3), or with recombinant baculovirus for P501 at different MOIs (lane 4 - 8). Cell lysates were run on SDS-PAGE under the reducing conditions and analyzed by Western blot with a monoclonal antibody against P5018 (P5018-10E3-G4D3). Lane 1 is the biotinylated protein molecular weight marker (StoLabs).



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Fig. 8

transmembrane, cytoplasmic, and extracellular regions Figure 1. Schematic of P501S with predicted

MYQRIWYSRLLRHRK AQLLLYNLLTFGLEYCLAAGIT YVPPI.LI.EVGVEEKFNI TMYLGIGPVI.GLYCVPLI.GSAS

DHWRGRYGRRRP FIWALSLOILLSLFLIPRAGWI, AGLI.CPDPRPI.E LALI ILGVOLL DFCGOVCFTPL

EALLSDLFRDPDHCRQ AYSYYAFMISLGGCLGYLLPAL DWIYISALAPYLGTQEE

CLEGILITLIELICYAATILY AEEAALGPTEPAEGISAPSISPHCCPCRARLAFRALGALLPRL

HOLCCRMPRTIAR LIVAGECSWMALMITTLEYTDE VOEGLYOGVPRARPGTEARRIYDEGVR

MOSIGLFLOCAISLYFSLVM DRLVQRFGTRAVYLAS YAAFPVAAGATCLSHSVAVVTA SAA

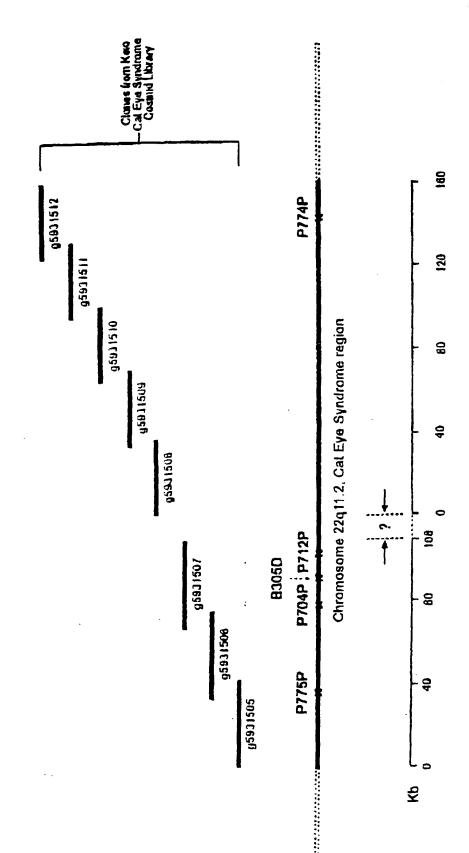
LTGETESALOILPYTLASLY HREKQVFLPKYRGDTGGASSEDSLMTSFLPGPKPGAPFPNGIIVGAGGSGL

LPPPALCGASACDVSVRVVVGEPTEARVVPGRG ICLIILAILDSAFLLSOVAPSLF MGSIVQLSQS

VTAYMVSAAGLGLVAIYFAT QVVFDKSDLAKYSA

Italic sequence: Predicted intracellular domain. Sequence in bold/underlined: used to generate polyclonal rabbit serum <u>Underlined sequence</u>: Predicted transmembrane domain; Bold sequence: Predicted extracellular domain;

Governing Amino Acid Composition of Integral Membrane Proteins: Applications to topology Prediction. J. Mol Biol. 283, Localization of domains predicted using HMMTOP (G.E. Tusnady and I. Simon (1998) Principles



Genomic Map of (5) Corixa Candidate Genes

Fig. 10

FIGURE 4. Elisa assay of rabbit polyclonal antibody specificity

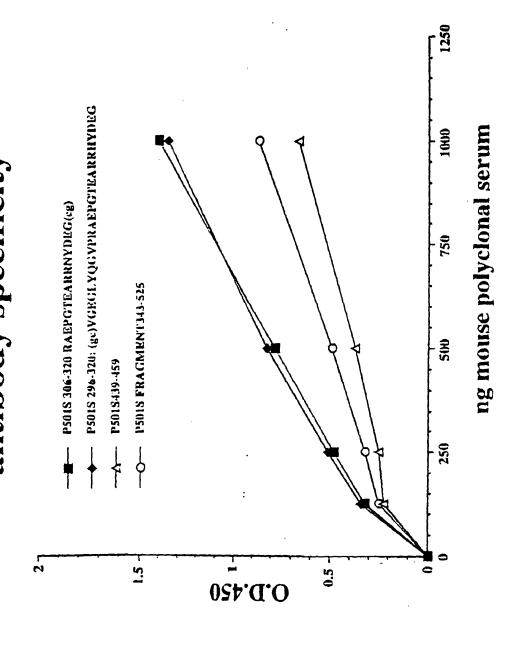


Fig. 11